STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10517079

Source: EFW

Date Processed by STIC: 11/28/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
        http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 10517,079C

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics
Wrapped Aminos

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 ____ Invalid Line Length

The rules require that a line **not exceed 72** characters in length. This includes white spaces.

3 ____ Misaligned Amino
Numbering

The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use **space characters**, instead.

4 ____ Non-ASCII

The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5 ____ Variable Length

Sequence(s) ____ contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ PatentIn 2.0
"bug"

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____ . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7 ____ Skipped Sequences
(OLD RULES)

Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8 ____ Skipped Sequences
(NEW RULES)

Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 ____ Use of n's or Xaa's
(NEW RULES)

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.

10 ____ Invalid <213>
Response

Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence

11 ____ Use of <220>

Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12 ____ PatentIn 2.0
"bug"

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 ____ Misuse of n/Xaa

"n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

IFWO

RAW SEQUENCE LISTING   DATE: 11/28/2006
PATENT APPLICATION: US/10/517,079C   TIME: 11:42:31

Input Set : **N:\efs\11_28_06**
\10517079c_efs\Substitute_Sequence_Listing_20747_230.txt
Output Set: **N:\CRF4\11282006\J517079C.raw**

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    4 <110> APPLICANT: CePeP AB
    6 <120> TITLE OF INVENTION: Cell Penetrating Peptides
    9 <130> FILE REFERENCE: 20747/230
   11 <140> CURRENT APPLICATION NUMBER: US 10/517,079C
C--> 12 <141> CURRENT FILING DATE: 2004-12-06
   14 <150> PRIOR APPLICATION NUMBER: PCT/IB03/03163
   15 <151> PRIOR FILING DATE: 2003-06-18
   17 <150> PRIOR APPLICATION NUMBER: US 60/391,788
   18 <151> PRIOR FILING DATE: 2002-06-25
   20 <150> PRIOR APPLICATION NUMBER: SE 0201863-8
   21 <151> PRIOR FILING DATE: 2002-06-18
   23 <160> NUMBER OF SEQ ID NOS: 31933
   25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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Does Not Comply
Corrected Diskette Needed

(pg. 1-6)

ERRORED SEQUENCES

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   38551 <210> SEQ ID NO: 2739
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   38553 <212> TYPE: PRT
   38554 <213> ORGANISM: Homo Sapiens
   38556 <220> FEATURE:
   38557 <222> LOCATION: ( 259 )...( 275 )
   38558 <223> OTHER INFORMATION: 5H6_HUMAN
W--> 38559 <400> SEQUENCE: 2739
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E--> 38561 1               5                   10                  15
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   38566 <211> LENGTH: 16
   38567 <212> TYPE: PRT
   38568 <213> ORGANISM: Homo Sapiens
   38570 <220> FEATURE:
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   38572 <223> OTHER INFORMATION: 5H6_HUMAN
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E--> 38575 1               5                   10                  15
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   38584 <220> FEATURE:
   38585 <222> LOCATION: ( 261 )...( 277 )
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See item #3 on = error summary sheet.

RAW SEQUENCE LISTING   DATE: 11/28/2006
PATENT APPLICATION: US/10/517,079C   TIME: 11:42:33

Input Set : N:\efs\11_28_06
\10517079c_efs\Substitute_Sequence_Listing_20747_230.txt
Output Set: N:\CRF4\11282006\J517079C.raw

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        38595 <212> TYPE: PRT
        38596 <213> ORGANISM: Homo Sapiens
        38598 <220> FEATURE:
        38599 <222> LOCATION: ( 108 )...( 124 )
        38600 <223> OTHER INFORMATION: 5H7_HUMAN
W--> 38601 <400> SEQUENCE: 2742
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E--> 38603 1               5               10              15
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        38608 <211> LENGTH: 16
        38609 <212> TYPE: PRT
        38610 <213> ORGANISM: Homo Sapiens
        38612 <220> FEATURE:
        38613 <222> LOCATION: ( 109 )...( 125 )
        38614 <223> OTHER INFORMATION: 5H7_HUMAN
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E--> 38617 1               5               10              15
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        38626 <220> FEATURE:
        38627 <222> LOCATION: ( 110 )...( 126 )
        38628 <223> OTHER INFORMATION: 5H7_HUMAN
W--> 38629 <400> SEQUENCE: 2744
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E--> 38631 1               5               10              15
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        38638 <213> ORGANISM: Homo Sapiens
        38640 <220> FEATURE:
        38641 <222> LOCATION: ( 305 )...( 321 )
        38642 <223> OTHER INFORMATION: 5H7_HUMAN
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        38652 <213> ORGANISM: Homo Sapiens
        38654 <220> FEATURE:
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Same errors ↓

See item #3 on error Summary Sheet

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                 RAW SEQUENCE LISTING              DATE: 11/28/2006
                 PATENT APPLICATION:  US/10/517,079C    TIME: 11:42:33

                 Input Set : N:\efs\11_28_06
\10517079c_efs\Substitute_Sequence_Listing_20747_230.txt
                 Output Set: N:\CRF4\11282006\J517079C.raw


       38655 <222> LOCATION: ( 248 )...( 264 )
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       38664 <211> LENGTH: 16
       38665 <212> TYPE: PRT
       38666 <213> ORGANISM: Homo Sapiens
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       38669 <222> LOCATION: ( 261 )...( 277 )
       38670 <223> OTHER INFORMATION: A1AA_HUMAN
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       38684 <223> OTHER INFORMATION: A1AA_HUMAN
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       38694 <213> ORGANISM: Homo Sapiens
       38696 <220> FEATURE:
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E--> 38715 1                 5                 10                15
       38719 <210> SEQ ID NO: 2751
       38720 <211> LENGTH: 16
       38721 <212> TYPE: PRT
       38722 <213> ORGANISM: Homo Sapiens
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SAme errors

See item #3 on error Summary Sheet.

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                    RAW SEQUENCE LISTING              DATE: 11/28/2006
                    PATENT APPLICATION:  US/10/517,079C    TIME: 11:42:33

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\10517079c_efs\Substitute_Sequence_Listing_20747_230.txt
                    Output Set: N:\CRF4\11282006\J517079C.raw


        38724 <220> FEATURE:
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        38726 <223> OTHER INFORMATION: A1AB_HUMAN
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E--> 38729 1               5               10              15
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        38734 <211> LENGTH: 16
        38735 <212> TYPE: PRT
        38736 <213> ORGANISM: Homo Sapiens
        38738 <220> FEATURE:
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        38740 <223> OTHER INFORMATION: A1AB_HUMAN
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        38752 <220> FEATURE:
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E--> 38785 1               5               10              15
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        38791 <212> TYPE: PRT
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Same errors (handwritten annotation)

RAW SEQUENCE LISTING   DATE: 11/28/2006
PATENT APPLICATION: US/10/517,079C   TIME: 11:42:33

Input Set : **N:\efs\11_28_06**
\10517079c_efs\Substitute_Sequence_Listing_20747_230.txt
Output Set: **N:\CRF4\11282006\J517079C.raw**

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E--> **38799** 1   5   10   15
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E--> **38813** 1   5   10   15
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W--> **38825 <400> SEQUENCE: 2758**
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E--> **38827** 1   5   10   15
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E--> **38855** 1   5   10   15
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          38860 <211> LENGTH: 16

[Handwritten notes:] Same errors

See item #3 on error summary sheet

RAW SEQUENCE LISTING   DATE: 11/28/2006
PATENT APPLICATION: US/10/517,079C   TIME: 11:42:33

Input Set : N:\efs\11_28_06
\10517079c_efs\Substitute_Sequence_Listing_20747_230.txt
Output Set: N:\CRF4\11282006\J517079C.raw

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E--> 38869 1               5                  10                  15
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      38874 <211> LENGTH: 16
      38875 <212> TYPE: PRT
      38876 <213> ORGANISM: Homo Sapiens
      38878 <220> FEATURE:
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      38894 <223> OTHER INFORMATION: A1AB_HUMAN
W--> 38895 <400> SEQUENCE: 2763
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      38904 <213> ORGANISM: Homo Sapiens
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      38917 <212> TYPE: PRT
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      38922 <223> OTHER INFORMATION: A1AD_HUMAN
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E--> 38925 1               5                  10                  15
      38929 <210> SEQ ID NO: 2766
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Same errors

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.